

Fig. 1

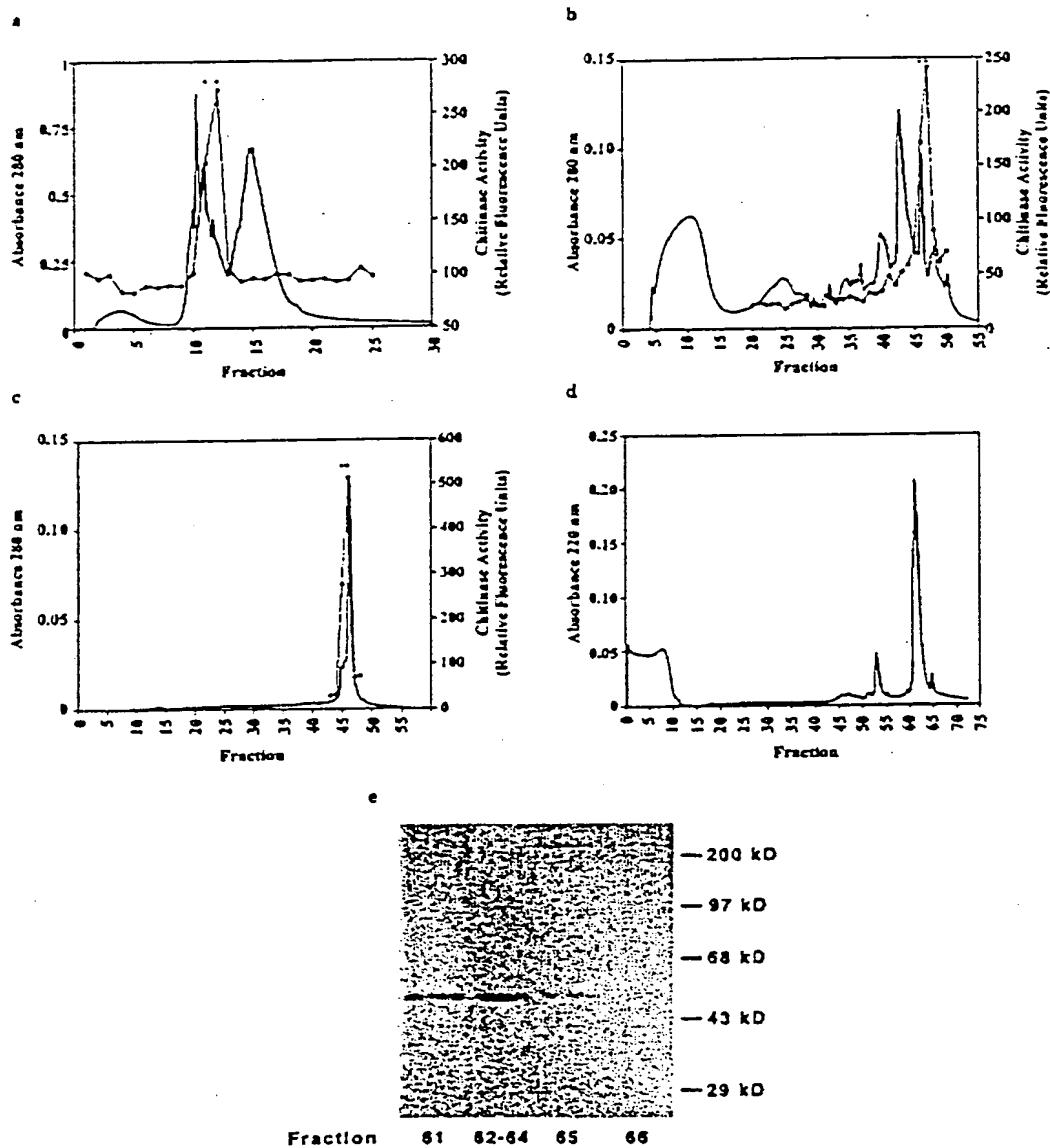


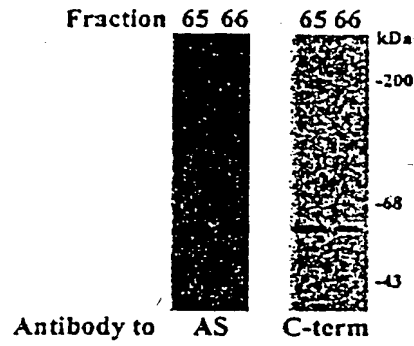
Diagram illustrating the 1.8 kb genomic region of the 5' end of the hTERT gene. The diagram shows a double-stranded DNA segment. The top strand (coding strand) has a 5' PCR primer binding site, followed by an ATG start codon, a GT29 insertion site, a Primer 2583 binding site, a GT34 insertion site, and a TAA stop codon. The bottom strand (template strand) has a Primer 2581 binding site and an oligo dT binding site. The distance between the 5' PCR primer and the TAA stop codon is indicated as 1.8 kb.

Fig. 3

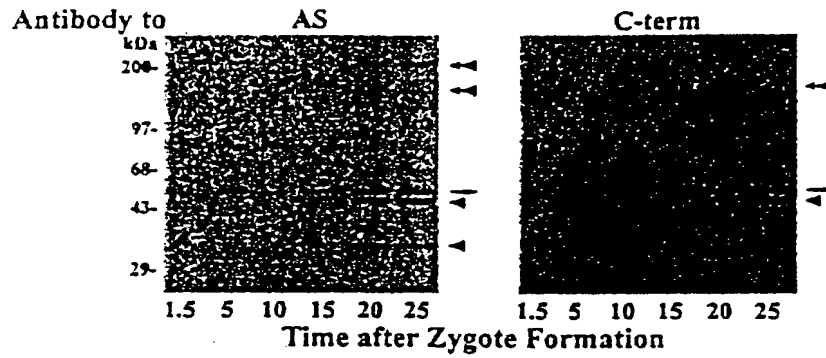
[illegible][illegible]

Fig. 4

a



b



c

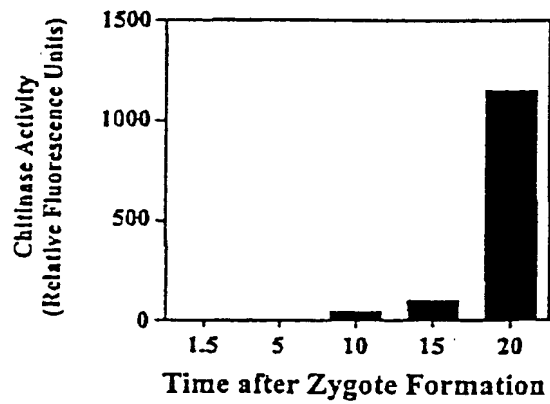


Fig. 5

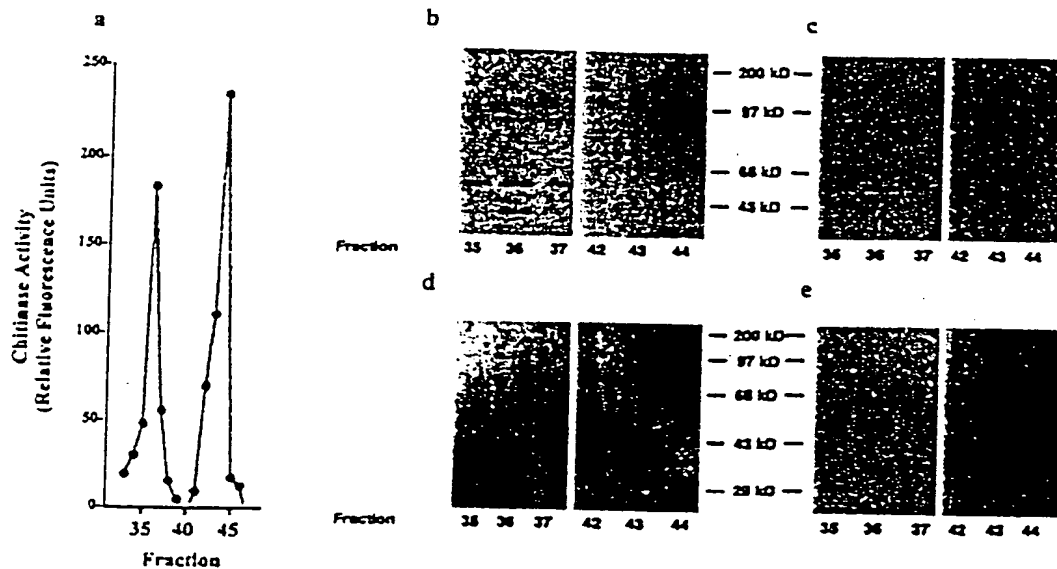


Fig. 6

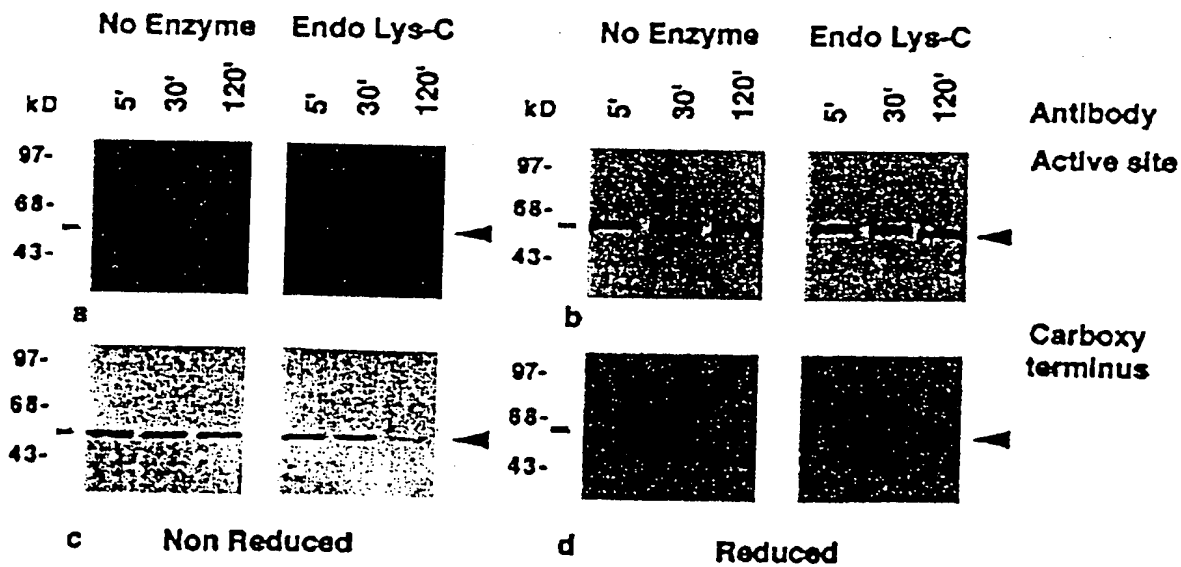


Fig. 7

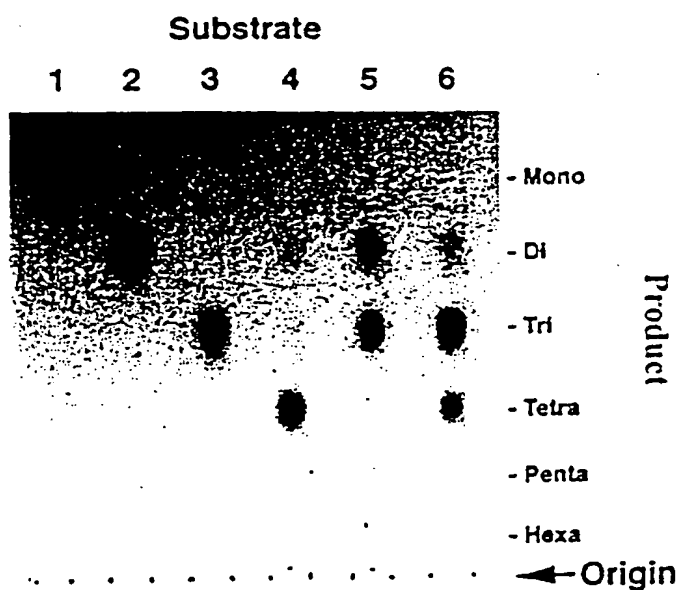
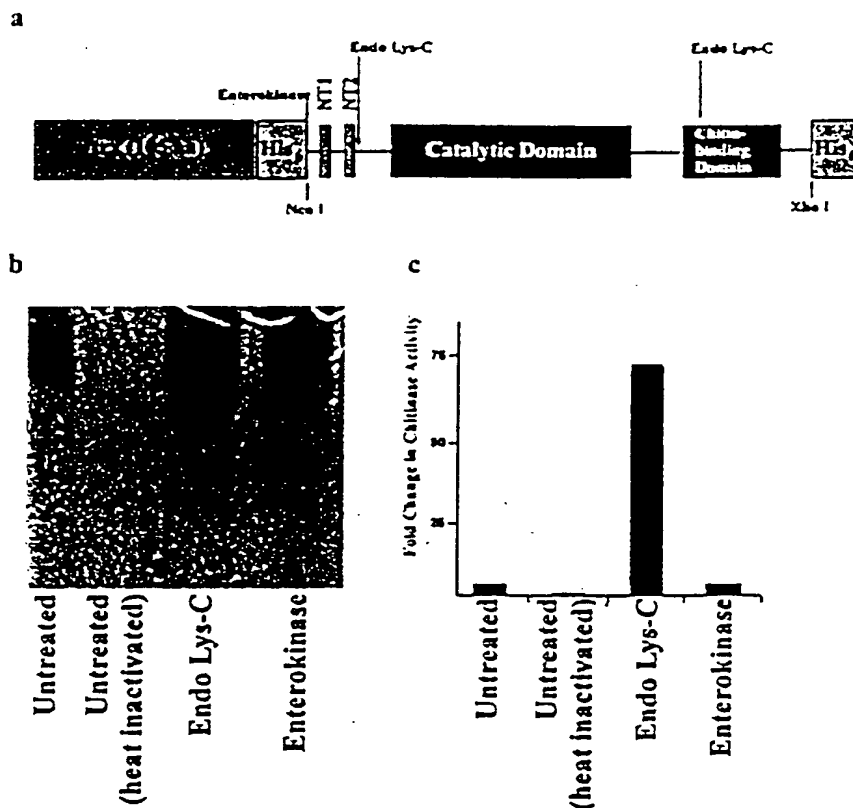
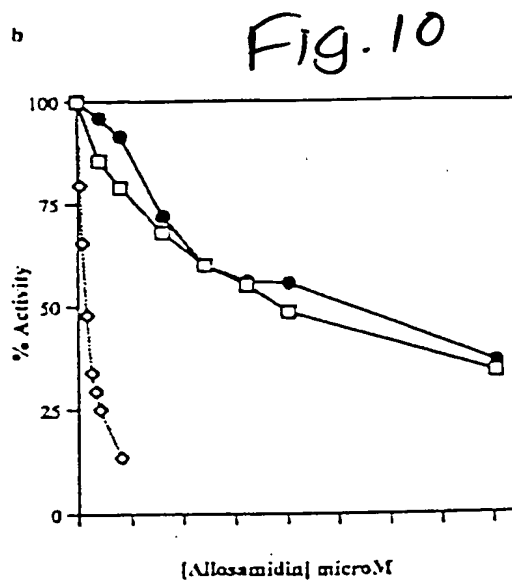
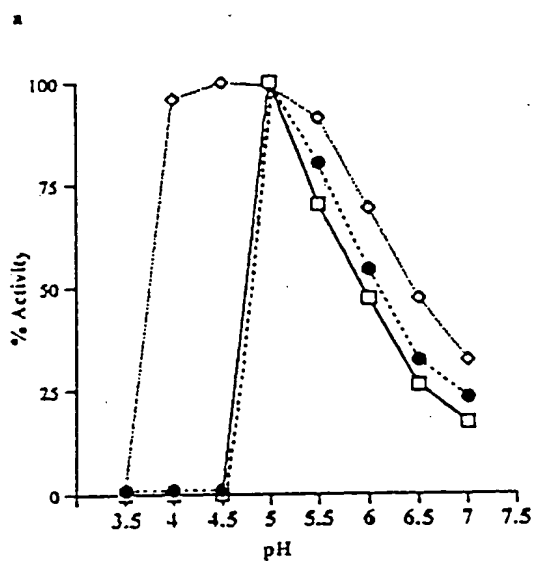
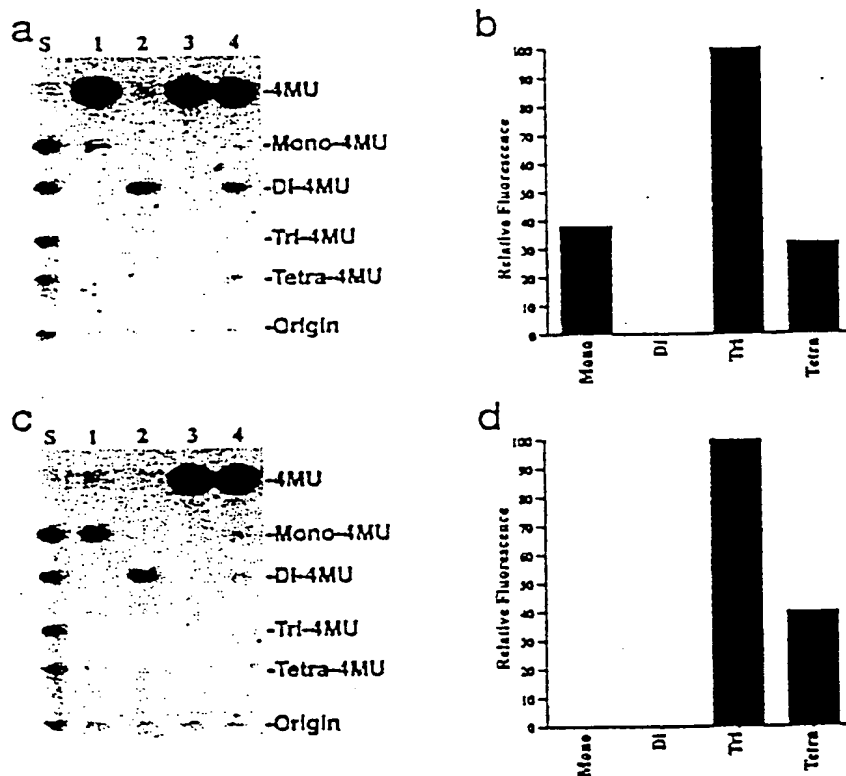


Fig. 8

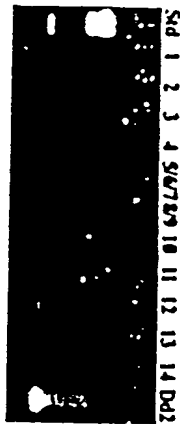
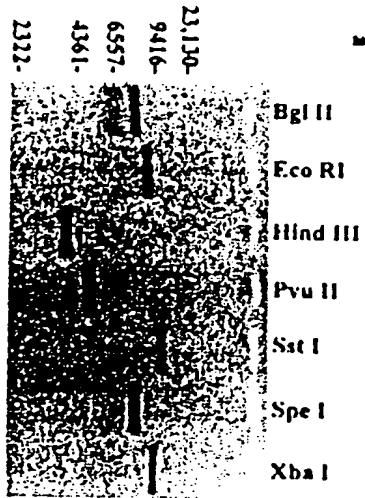
Fig. 9



[illegible]

0609

Fig. 12



09579383.052600

Fig. 13

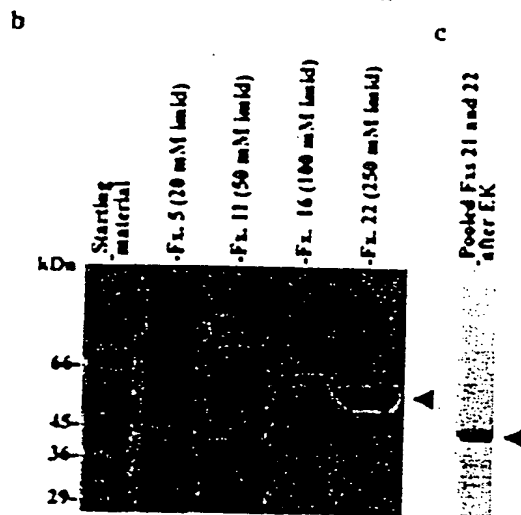
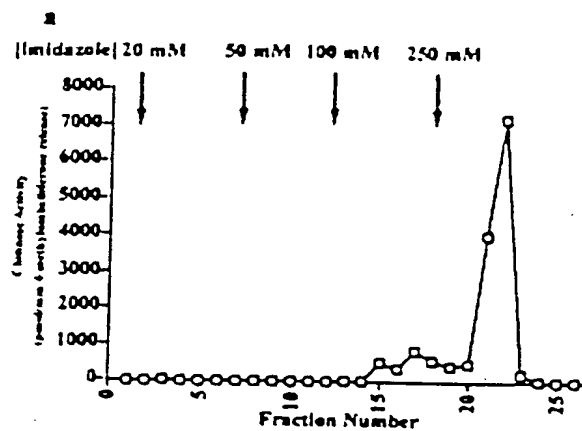
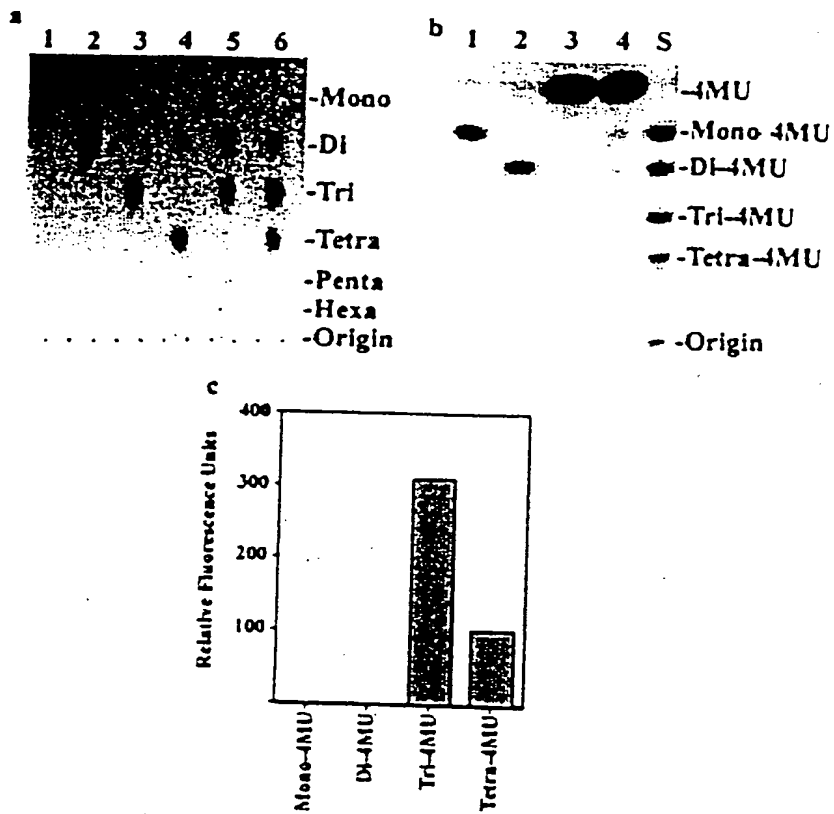


Fig. 14



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Fig. 15

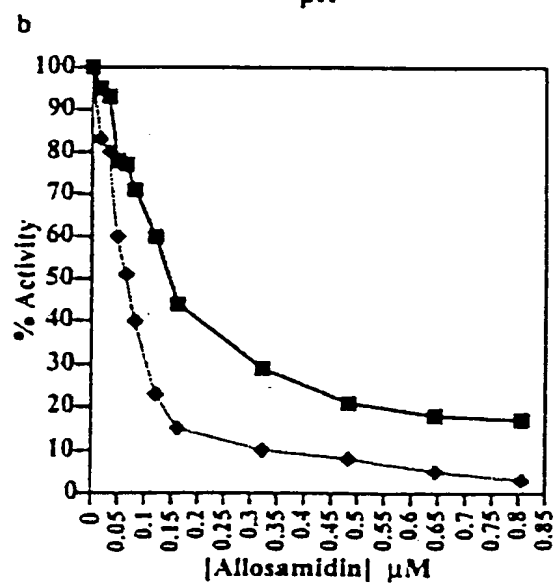
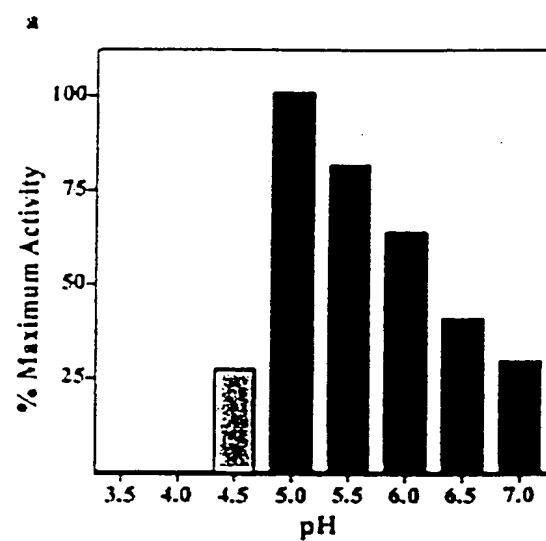


Fig. 16



009250*E8E6/560